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================================================================
Sequence Listing could not be accepted.
If you need help call the Patent Electronic Business Center at (866)
217-9197 (toll free).
Reviewer: Anne Corrigan
Timestamp: [year=2008; month=3; day=18; hr=17; min=45; sec=55; ms=198; ]
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****************************************************************
Reviewer Comments:
<210>   3
<211>   30
<212>   DNA
<213>   Sense primer

<400>   3
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The above <213> response is invalid, per 1.823 of the Sequence Rules.
The only valid responses are:  the Genus species of the organism,
"Artificial Sequence," or "Unknown."  "Artificial Sequence" and
"Unknown" require explanation in the <220>-<223> section.  Same error in
Sequences 4-6.
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Application No: 10590034   Version No: 1.0

Input Set:

Output Set:

Started: 2008-03-06 15:23:15.961
Finished: 2008-03-06 15:23:16.241
Elapsed: 0 hr(s) 0 min(s) 0 sec(s) 280 ms
Total Warnings: 4
Total Errors: 0
No. of SeqIDs Defined: 6
Actual SeqID Count: 6

Error code   Error Description

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W   402   Undefined organism found in <213> in SEQ ID (6)

<110>   Genomine, Inc.
        Korea Research Institute of Chemical Technology

<120>   Polypeptide Participating in Pyridoxine Biosynthesis, a
        Polynucleotide Coding the Polypeptide and Those Uses

<130>   DJKIM.GENO.PT1

<140>   10590034
<141>   2008-03-06

<150>   PCT/KR05/000453
<151>   2006-08-18

<150>   PCT/KR2005/000453
<151>   2005-02-18

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cggcgttgtg gcggtgtacg gtaacggtgc gataacggag gcgaagaaat ctcccttctc   180

cgtgaaggtc ggtttggctc agatgctccg tggtggtgtt atcatggatg tcgtcaacgc   240

cgagcaagct cgtatcgccg aggaggctgg tgcttgcgcc gtcatggctt ggagcgtgt   300

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        35   40   45

Ile Ala Glu Glu Ala Gly Ala Cys Ala Val Met Ala Leu Glu Arg Val
    50   55   60

Pro Ala Asp Ile Arg Ala Gln Gly Gly Val Ala Arg Met Ser Asp Pro
65   70   75   80

Gln Met Ile Lys Glu Ile Lys Gln Ala Val Thr Ile Pro Val Met Ala
                85   90   95

Lys Ala Arg Ile Gly His Phe Val Glu Ala Gln Ile Leu Glu Ala Ile
                100   105   110

Gly Ile Asp Tyr Ile Asp Glu Ser Glu Val Leu Thr Leu Ala Asp Glu
        115   120   125

Asp His His Ile Asn Lys His Asn Phe Arg Ile Pro Phe Val Cys Gly
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Cys Arg Asn Leu Gly Glu Ala Leu Arg Arg Ile Arg Glu Gly Ala Ala
145   150   155   160

Met Ile Arg Thr Lys Gly Glu Ala Gly Thr Gly Asn Ile Ile Glu Ala
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Val Arg His Val Arg Ser Val Asn Gly Asp Ile Arg Val Leu Arg Asn
             180   185   190

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Tyr Asp Leu Val Met Gln Thr Lys Gln Leu Gly Arg Leu Pro Val Val
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Gln Phe Ala Ala Gly Gly Val Ala Thr Pro Ala Asp Ala Ala Leu Met
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His Tyr Ser Asp Pro Glu Met Leu Val Glu Val Ser Cys Gly Leu Gly
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Ala Asn Arg Ser Glu
305

<210>   3
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<212>   DNA
<213>   Sense primer

<400>   3
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<210> 4
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